

KEY_Practice24_Basic_Stats_IV_Significance

December 10, 2021

1 Practice: Statistical Significance

Let's continue to work with the diabetes dataset to apply a t-test to real world data.

```
[1]: # Import pandas, so that we can import the diabetes dataset and work with the
      ↪data frame version of this data
import pandas as pd
```

```
[2]: # Set the path
path = 'https://raw.githubusercontent.com/GWC-DCMB/curriculum-notebooks/master/'
# This is where the file is located
filename = path + 'SampleData/diabetes.csv'
```

```
[3]: # Load the diabetes dataset into a DataFrame
diabetes_df = pd.read_csv(filename)
diabetes_df
```

```
[3]:
```

	AGE	SEX	BMI	MAP	TC	LDL	HDL	TCH	LTG	GLU	Y
0	59	2	32.1	101.00	157	93.2	38.0	4.00	4.8598	87	151
1	48	1	21.6	87.00	183	103.2	70.0	3.00	3.8918	69	75
2	72	2	30.5	93.00	156	93.6	41.0	4.00	4.6728	85	141
3	24	1	25.3	84.00	198	131.4	40.0	5.00	4.8903	89	206
4	50	1	23.0	101.00	192	125.4	52.0	4.00	4.2905	80	135
...
437	60	2	28.2	112.00	185	113.8	42.0	4.00	4.9836	93	178
438	47	2	24.9	75.00	225	166.0	42.0	5.00	4.4427	102	104
439	60	2	24.9	99.67	162	106.6	43.0	3.77	4.1271	95	132
440	36	1	30.0	95.00	201	125.2	42.0	4.79	5.1299	85	220
441	36	1	19.6	71.00	250	133.2	97.0	3.00	4.5951	92	57

[442 rows x 11 columns]

We are interested in understanding whether there are differences in LDL levels (the “bad” cholesterol) by sex, i.e. are LDL levels different for males vs. females?

1. Formulate the null hypothesis and the alternative hypothesis. - **Null hypothesis:** There is NO difference in LDL levels between male and female. - **Alternative hypothesis:** There is a difference in LDL levels by sex.

```
[4]: # Import numpy
import numpy as np
```

Males are indicated by “1” for the variable “SEX”, while females are indicated by “2”.

```
[5]: # Define a vector of the LDL levels for males and name it ldl_male
diabetes_male = diabetes_df.query('SEX == 1')
ldl_male = diabetes_male['LDL']

# Define a vector of the LDL levels for females and name it ldl_female
diabetes_female = diabetes_df.query('SEX == 2')
ldl_female = diabetes_female['LDL']
```

2. Identify and compute a test statistic that can be used to reject or fail to reject the null hypothesis. - As we are working with two independent samples, we will use the two-sample t-test and use the t-statistic.

3. Compute the test statistic and p-value.

```
[6]: # Import stats methods to help calculate the t-statistic and p-value
from scipy import stats
```

```
[7]: # Run a Student's t-test
t_statistic, p_value = stats.ttest_ind(ldl_male, ldl_female)

# Print out the test statistic and p-value
print("t-statistic = " + str(t_statistic))
print("p-value = " + str(p_value))
```

t-statistic = -3.022893334345971

p-value = 0.0026499873735660695

4. Compare the p-value to an acceptable significance value, α and compare the test statistic to acceptable critical value(s). If $p\text{-value} \leq \alpha$ and the test-statistic $\geq +\text{critical value}$ or test-statistic $\leq -\text{critical value}$, that the observed effect is statistically significant, the null hypothesis is rejected, and the alternative hypothesis is valid.** - $p\text{-value} = 0.0026 < 0.05$, so we reject the null hypothesis. - $t\text{-statistic} = -3.02 < -1.96$, so this reaffirms that we reject the null hypothesis. - Interpretation: There is a significant difference in LDL levels between males and females.

Congratulations on completing the lesson and practice!

It's a lot of information, but you learned powerful tools to be on your way to answer your own research questions by analyzing real world data!

Challenge: Using the code you wrote above as a template, can you run a t-test comparing LDL Cholesterol for people 50 & older vs. people under 50?

```
[8]: # Define a vector of the LDL levels for people 50 or older
diabetes_over50 = diabetes_df.query('AGE >= 50')
```

```
ldl_over50 = diabetes_over50['LDL']

# Define a vector of the LDL levels for females and name it ldl_female
diabetes_under50 = diabetes_df.query('AGE < 50')
ldl_under50 = diabetes_under50['LDL']

# Run a Student's t-test
t_statistic, p_value = stats.ttest_ind(ldl_over50, ldl_under50)

# Print out the test statistic and p-value
print("t-statistic = " + str(t_statistic))
print("p-value = " + str(p_value))
```

```
t-statistic = 3.185760417933572
p-value = 0.001546465356577734
```